

0590
0327

#2

OIPe

RAW SEQUENCE LISTING

DATE: 04/01/2002

PATENT APPLICATION: US/10/036,434

TIME: 14:52:14

Input Set : N:\Crf3\RULE60\10036434.raw

Output Set: N:\CRF3\04012002\J036434.raw

1 <110> APPLICANT: WALLACH, David
 2 ENGELMANN, Hartmut
 3 ADERKA, Dan
 4 RUBINSTEIN, Menachem
 5 <120> TITLE OF INVENTION: TUMOR NECROSIS FACTOR INHIBITORY PROTEIN AND ITS
 PURIFICATION
 6 <130> FILE REFERENCE: WALLACH1C
 7 <140> CURRENT APPLICATION NUMBER: 10/036,434
 8 <141> CURRENT FILING DATE: 2002-01-07
 10 <150> PRIOR APPLICATION NUMBER: 09/414,609
 11 <151> PRIOR FILING DATE: 1999-10-08
 14 <150> PRIOR APPLICATION NUMBER: 08/474,691
 15 <151> PRIOR FILING DATE: 1995-06-07
 16 <150> PRIOR APPLICATION NUMBER: 07/876,828
 17 <151> PRIOR FILING DATE: 1992-04-30
 18 <150> PRIOR APPLICATION NUMBER: 07/243,092
 19 <151> PRIOR FILING DATE: 1988-09-12
 20 <160> NUMBER OF SEQ ID NOS: 1
 21 <170> SOFTWARE: PatentIn version 3.0
 23 <210> SEQ ID NO: 1
 24 <211> LENGTH: 16
 25 <212> TYPE: PRT
 26 <213> ORGANISM: Artificial Sequence
 27 <220> FEATURE:
 28 <223> OTHER INFORMATION: synthetic
 W--> 29 <221> NAME/KEY: misc_feature
 30 <222> LOCATION: (14)..(14)
 31 <223> OTHER INFORMATION: Xaa is unknown
 W--> 32 <400> 1
 W--> 33 Asp Ser Val Cys Pro Gln Gly Lys Tyr Ile His Pro Gln Xaa Asn Ser
 34 1 5 10 15

ENTERED

RAW SEQUENCE LISTING ERROR SUMMARY
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Output Set: N:\CRF3\04012002\J036434.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:1; Xaa Pos. 14

Invalid Line Length:

The rules require that a line not exceed 72 characters in length. This includes spaces.

Seq#:1; Line(s) 5

VERIFICATION SUMMARY

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Input Set : N:\Crf3\RULE60\10036434.raw

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L:29 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!

L:32 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:1

L:33 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:0